

SEQUENCE LISTING

<110> DENEFLÉ, Patrice
 ROSIER-MONTUS, Marie-Francoise
 PRADES, Catherine
 ARNOULD-REGUIGNE, Isabelle
 DUVERGER, Nicolas
 ALLIKMETS, Rando
 DEAN, Michael

<120> NUCLEIC ACIDS OF THE HUMAN ABCA5, ABCA6, ABCA9, AND ABCA10 GENES, VECTORS
 CONTAINING SUCH NUCLEIC ACIDS, AND USES THEREOF

<130> ABCA5, 6, 9, 10

<140> US 10/005,338

<141> 2001-12-07

<150> US 60/263,231

<151> 2001-01-23

<150> FR 00403440.1

<151> 2000-12-07

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<170> PatentIn Ver. 2.1

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gtaagaaagt accgcagtta atattttctt tagccaactt atattcaatg tattttttat 6120
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Val Gln Glu Ile Leu Phe Pro Leu Phe Phe Leu Phe Trp Leu Ile Leu
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Ile Ser Met Met His Pro Asn Lys Lys Tyr Glu Glu Val Pro Asn Ile
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Glu Leu Asn Pro Met Asp Lys Phe Thr Leu Ser Asn Leu Ile Leu Gly
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Tyr Thr Pro Val Thr Asn Ile Thr Ser Ser Ile Met Gln Lys Val Ser
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Thr Asp His Leu Pro Asp Val Ile Ile Thr Glu Glu Tyr Thr Asn Glu
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Lys Glu Met Leu Thr Ser Ser Leu Ser Lys Pro Ser Asn Phe Val Gly
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Val Val Phe Lys Asp Ser Met Ser Tyr Glu Leu Arg Phe Phe Pro Asp
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Met Ile Pro Val Ser Ser Ile Tyr Met Asp Ser Arg Ala Gly Cys Ser

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Phe	Lys	Lys	Ser	Lys	His	Val	Gly	Ile	Val	Glu	Phe	Phe	Val	Thr	Val
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Ile	Met	Leu	Thr	Leu	Asn	Ser	Ile	Phe	Tyr	Val	Leu	Leu	Ala	Val	Tyr
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Tyr	Phe	Leu	Lys	Pro	Ser	Tyr	Trp	Ser	Lys	Ser	Lys	Arg	Asn	Tyr	Glu
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 Leu Gly His Ser Gly Thr Gly Lys Ser Thr Leu Met Asn Ile Leu Cys
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 Gly Leu Cys Pro Pro Ser Asp Gly Phe Ala Ser Ile Tyr Gly His Arg
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 Val Ser Glu Ile Asp Glu Met Phe Glu Ala Arg Lys Met Ile Gly Ile
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 Cys Pro Gln Leu Asp Ile His Phe Asp Val Leu Thr Val Glu Glu Asn
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 Leu Ser Ile Leu Ala Ser Ile Lys Gly Ile Pro Ala Asn Asn Ile Ile
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 Gln Glu Val Gln Lys Val Leu Leu Asp Leu Asp Met Gln Thr Ile Lys
 595 600 605
 Asp Asn Gln Ala Lys Lys Leu Ser Gly Gly Gln Lys Arg Lys Leu Ser
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 Leu Gly Ile Ala Val Leu Gly Asn Pro Lys Ile Leu Leu Leu Asp Glu
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 Pro Thr Ala Gly Met Asp Pro Cys Ser Arg His Ile Val Trp Asn Leu
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 Leu Lys Tyr Arg Lys Ala Asn Arg Val Thr Val Phe Ser Thr His Phe
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 Met Asp Glu Ala Asp Ile Leu Ala Asp Arg Lys Ala Val Ile Ser Gln
 675 680 685
 Gly Met Leu Lys Cys Val Gly Ser Ser Met Phe Leu Lys Ser Lys Trp
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 Gly Ile Gly Tyr Arg Leu Ser Met Tyr Ile Asp Lys Tyr Cys Ala Thr
 705 710 715 720
 Glu Ser Leu Ser Ser Leu Val Lys Gln His Ile Pro Gly Ala Thr Leu
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 Leu Gln Gln Asn Asp Gln Gln Leu Val Tyr Ser Leu Pro Phe Lys Asp
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 Phe Leu Lys Leu Glu Val Glu Ala Glu Ile Asp Gln Ala Asp Tyr Ser
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 Val Phe Thr Gln Gln Pro Leu Glu Glu Glu Met Asp Ser Lys Ser Phe
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Leu Val Ser Thr Met Ser Leu Trp Lys Gln Gln Met Tyr Thr Ile Ala
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 Lys Phe His Phe Phe Thr Leu Lys Arg Glu Ser Lys Ser Val Arg Ser
 850 855 860
 Val Leu Leu Leu Leu Leu Ile Phe Phe Thr Val Gln Ile Phe Met Phe
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 Leu Val His His Ser Phe Lys Asn Ala Val Val Pro Ile Lys Leu Val
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 Pro Asp Leu Tyr Phe Leu Lys Pro Gly Asp Lys Pro His Lys Tyr Lys
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 Thr Ser Leu Leu Leu Gln Asn Ser Ala Asp Ser Asp Ile Ser Asp Leu
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 Ile Ser Phe Phe Thr Ser Gln Asn Ile Met Val Thr Met Ile Asn Asp
 930 935 940
 Ser Asp Tyr Val Ser Val Ala Pro His Ser Ala Ala Leu Asn Val Met
 945 950 955 960
 His Ser Glu Lys Asp Tyr Val Phe Ala Ala Val Phe Asn Ser Thr Met
 965 970 975
 Val Tyr Ser Leu Pro Ile Leu Val Asn Ile Ile Ser Asn Tyr Tyr Leu
 980 985 990
 Tyr His Leu Asn Val Thr Glu Thr Ile Gln Ile Trp Ser Thr Pro Phe
 995 1000 1005
 Phe Gln Glu Ile Thr Asp Ile Val Phe Lys Ile Glu Leu Tyr Phe Gln
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 Ala Ala Leu Leu Gly Ile Ile Val Thr Ala Met Pro Pro Tyr Phe Ala
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 Met Glu Asn Ala Glu Asn His Lys Ile Lys Ala Tyr Thr Gln Leu Lys
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 1060 1065 1070
 Asp Ile Pro Leu Phe Phe Ile Ile Leu Ile Leu Met Leu Gly Ser Leu
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 Val Val Phe Cys Leu Ile Gly Tyr Val Pro Ser Val Ile Leu Phe Thr
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 Tyr Ile Ala Ser Phe Thr Phe Lys Lys Ile Leu Asn Thr Lys Glu Phe
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 Trp Ser Phe Ile Tyr Ser Val Ala Ala Leu Xaa Cys Ile Ala Ile Thr
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 Glu Ile Thr Phe Phe Met Gly Tyr Thr Ile Ala Thr Ile Leu His Tyr
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 Ala Phe Cys Ile Ile Ile Pro Ile Tyr Pro Leu Leu Gly Cys Leu Ile

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Ser Phe Ile Lys Ile Ser Trp Lys Asn Val Arg Lys Asn Val Asp Thr		
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Tyr Asn Pro Trp Asp Arg Leu Ser Val Ala Val Ile Ser Pro Tyr Leu		
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Gln Cys Val Leu Trp Ile Phe Leu Leu Gln Tyr Tyr Glu Lys Lys Tyr		
	1220	1225 1230
Gly Gly Arg Ser Ile Arg Lys Asp Pro Phe Phe Arg Asn Leu Ser Thr		
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Lys Ser Lys Asn Arg Lys Leu Pro Glu Pro Pro Asp Asn Glu Asp Glu		
	1250	1255 1260
Asp Glu Asp Val Lys Ala Glu Arg Leu Lys Val Lys Glu Leu Met Gly		
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Cys Gln Cys Cys Glu Glu Lys Pro Ser Ile Met Val Ser Asn Leu His		
	1285	1290 1295
Lys Glu Tyr Asp Asp Lys Lys Asp Phe Leu Leu Ser Arg Lys Val Lys		
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Lys Val Ala Thr Lys Tyr Ile Ser Phe Cys Val Lys Lys Gly Glu Ile		
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Leu Gly Leu Leu Gly Pro Asn Gly Ala Gly Lys Ser Thr Ile Ile Asn		
	1330	1335 1340
Ile Leu Val Gly Asp Ile Glu Pro Thr Ser Gly Gln Val Phe Leu Gly		
1345	1350	1355 1360
Asp Tyr Ser Ser Glu Thr Ser Glu Asp Asp Asp Ser Leu Lys Cys Met		
	1365	1370 1375
Gly Tyr Cys Pro Gln Ile Asn Pro Leu Trp Pro Asp Thr Thr Leu Gln		
	1380	1385 1390
Glu His Phe Glu Ile Tyr Gly Ala Val Lys Gly Met Ser Ala Ser Asp		
	1395	1400 1405
Met Lys Glu Val Ile Ser Arg Ile Thr His Ala Leu Asp Leu Lys Glu		
	1410	1415 1420
His Leu Gln Lys Thr Val Lys Lys Leu Pro Ala Gly Ile Lys Arg Lys		
1425	1430	1435 1440
Leu Cys Phe Ala Leu Ser Met Leu Gly Asn Pro Gln Ile Thr Leu Leu		
	1445	1450 1455
Asp Glu Pro Ser Thr Gly Met Asp Pro Lys Ala Lys Gln His Met Trp		
	1460	1465 1470
Arg Ala Ile Arg Thr Ala Phe Lys Asn Arg Lys Arg Ala Ala Ile Leu		
	1475	1480 1485
Thr Thr His Tyr Met Glu Glu Ala Glu Ala Val Cys Asp Arg Val Ala		
	1490	1495 1500
Ile Met Val Ser Gly Gln Leu Arg Cys Ile Gly Thr Val Gln His Leu		
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Lys Ser Lys Phe Gly Lys Gly Tyr Phe Leu Glu Ile Lys Leu Lys Asp
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 Trp Ile Glu Asn Leu Glu Val Asp Arg Leu Gln Arg Glu Ile Gln Tyr
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 Ile Phe Pro Asn Ala Ser Arg Gln Glu Ser Phe Ser Ser Ile Leu Ala
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 Tyr Lys Ile Pro Lys Glu Asp Val Gln Ser Leu Ser Gln Ser Phe Phe
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 Lys Leu Glu Glu Ala Lys His Ala Phe Ala Ile Glu Glu Tyr Ser Phe
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 Ser Gln Ala Thr Leu Glu Gln Val Phe Val Glu Leu Thr Lys Glu Gln
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 Ser Ser Met Arg Asn Val Gln Phe Pro Gly Met Ala Pro Gln Asn Leu
 50 55 60
 Gly Arg Val Asp Lys Phe Asn Ser Ser Ser Leu Met Val Val Tyr Thr
 65 70 75 80
 Pro Ile Ser Asn Leu Thr Gln Gln Ile Met Asn Lys Thr Ala Leu Ala
 85 90 95
 Pro Leu Leu Lys Gly Thr Ser Val Ile Gly Ala Pro Asn Lys Thr His
 100 105 110
 Met Asp Glu Ile Leu Leu Glu Asn Leu Pro Tyr Ala Met Gly Ile Ile
 115 120 125
 Phe Asn Glu Thr Phe Ser Tyr Lys Leu Ile Phe Phe Gln Gly Tyr Asn
 130 135 140
 Ser Pro Leu Trp Lys Glu Asp Phe Ser Ala His Cys Trp Asp Gly Tyr
 145 150 155 160
 Gly Glu Phe Ser Cys Thr Leu Thr Lys Tyr Trp Asn Arg Gly Phe Val

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Ala	Leu	Gln	Thr	Ala	Ile	Asn	Thr	Ala	Ile	Ile	Glu	Ile	Thr	Thr	Asn
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His	Pro	Val	Met	Glu	Glu	Leu	Met	Ser	Val	Thr	Ala	Ile	Thr	Met	Lys
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Thr	Leu	Pro	Phe	Ile	Thr	Lys	Asn	Leu	Leu	His	Asn	Glu	Met	Phe	Ile
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Leu	Phe	Phe	Leu	Leu	His	Phe	Ser	Pro	Leu	Val	Tyr	Phe	Ile	Ser	Leu
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Asn	Val	Thr	Lys	Glu	Arg	Lys	Lys	Ser	Lys	Asn	Leu	Met	Lys	Met	Met
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Gly	Leu	Gln	Asp	Ser	Ala	Phe	Trp	Leu	Ser	Trp	Gly	Leu	Ile	Tyr	Ala
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Gly	Phe	Ile	Phe	Ile	Ile	Ser	Ile	Phe	Ile	Thr	Ile	Ile	Ile	Thr	Phe
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Phe	Leu	Tyr	Gly	Leu	Ser	Leu	Val	Ala	Leu	Val	Phe	Leu	Met	Ser	Val
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Leu	Leu	Lys	Lys	Ala	Val	Leu	Thr	Asn	Leu	Val	Val	Phe	Leu	Leu	Thr
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Thr	Gly	Met	Ile	Gln	Ile	Ile	Lys	Leu	Asp	Tyr	Asn	Leu	Asn	Gly	Val
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Ser	Met	Leu	Leu	Leu	Asp	Gly	Leu	Ile	Tyr	Leu	Leu	Leu	Ala	Leu	Tyr
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Phe	Asp	Lys	Ile	Leu	Pro	Tyr	Gly	Asp	Glu	Arg	His	Tyr	Ser	Pro	Leu
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Phe	Phe	Leu	Asn	Ser	Ser	Ser	Cys	Phe	Gln	His	Gln	Arg	Thr	Asn	Ala
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Lys	Val	Ile	Glu	Lys	Glu	Ile	Asp	Ala	Glu	His	Pro	Ser	Asp	Asp	Tyr
	450					455					460				
Phe	Glu	Pro	Val	Ala	Pro	Glu	Phe	Gln	Gly	Lys	Glu	Ala	Ile	Arg	Ile
465					470					475					480
Arg	Asn	Val	Lys	Lys	Glu	Tyr	Lys	Gly	Lys	Ser	Gly	Lys	Val	Glu	Ala
				485					490					495	
Leu	Lys	Gly	Leu	Leu	Phe	Asp	Ile	Tyr	Glu	Gly	Gln	Ile	Thr	Ala	Ile
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 Gly Leu Ser Val Pro Thr Glu Gly Ser Val Thr Ile Tyr Asn Lys Asn
 530 535 540
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 545 550 555
 Cys Pro Gln Phe Asn Val Gln Phe Asp Ile Leu Thr Val Lys Glu Asn
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 Leu Ser Leu Phe Ala Lys Ile Lys Gly Ile His Leu Lys Glu Val Glu
 580 585 590
 Gln Glu Val Gln Arg Ile Leu Leu Glu Leu Asp Met Gln Asn Ile Gln
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 Asp Asn Leu Ala Lys His Leu Ser Glu Gly Gln Lys Arg Lys Leu Thr
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 Phe Gly Ile Thr Ile Leu Gly Asp Pro Gln Ile Leu Leu Leu Asp Glu
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 Pro Thr Thr Gly Leu Asp Pro Phe Ser Arg Asp Gln Val Trp Ser Leu
 645 650 655
 Leu Arg Glu Arg Arg Ala Asp His Val Ile Leu Phe Ser Thr Gln Ser
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 Met Asp Glu Ala Asp Ile Leu Ala Asp Arg Lys Val Ile Met Ser Asn
 675 680 685
 Gly Arg Leu Lys Cys Ala Gly Ser Ser Met Phe Leu Lys Arg Arg Trp
 690 695 700
 Gly Leu Gly Tyr His Leu Ser Leu His Arg Asn Glu Ile Cys Asn Pro
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 Glu Gln Ile Thr Ser Phe Ile Thr His His Ile Pro Asp Ala Lys Leu
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 Lys Thr Glu Asn Lys Glu Lys Leu Val Tyr Thr Leu Pro Leu Glu Arg
 740 745 750
 Thr Asn Thr Phe Pro Asp Leu Phe Ser Asp Leu Asp Lys Cys Ser Asp
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 Gln Gly Val Thr Gly Tyr Asp Ile Ser Met Ser Thr Leu Asn Glu Val
 770 775 780
 Phe Met Lys Leu Glu Gly Gln Ser Thr Ile Glu Gln Asp Phe Glu Gln
 785 790 795 800
 Val Glu Met Ile Arg Asp Ser Glu Ser Leu Asn Glu Met Glu Leu Ala
 805 810 815
 His Ser Ser Phe Ser Glu Met Gln Thr Ala Val Ser Asp Met Gly Leu
 820 825 830
 Trp Arg Met Gln Val Phe Ala Met Ala Arg Leu Arg Phe Leu Lys Leu
 835 840 845

Lys Arg Gln Thr Lys Val Leu Leu Thr Leu Leu Leu Val Phe Gly Ile
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 Ala Ile Phe Pro Leu Ile Val Glu Asn Ile Ile Tyr Ala Met Leu Asn
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 Glu Lys Ile Asp Trp Glu Phe Lys Asn Glu Leu Tyr Phe Leu Ser Pro
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 Gly Gln Leu Pro Gln Glu Pro Arg Thr Ser Leu Leu Ile Ile Asn Asn
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 Thr Glu Ser Asn Ile Glu Asp Phe Ile Lys Ser Leu Lys His Gln Asn
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 Ile Leu Leu Glu Val Asp Asp Phe Glu Asn Arg Asn Gly Thr Asp Gly
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 Leu Ser Tyr Asn Gly Ala Ile Ile Val Ser Gly Lys Gln Lys Asp Tyr
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 Arg Phe Ser Val Val Cys Asn Thr Lys Arg Leu His Cys Phe Pro Ile
 965 970 975
 Leu Met Asn Ile Ile Ser Asn Gly Leu Leu Gln Met Phe Asn His Thr
 980 985 990
 Gln His Ile Arg Ile Glu Ser Ser Pro Phe Pro Leu Ser His Ile Gly
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 Lys Asn Ala Lys Ser Gln Leu Trp Ile Ser Gly Leu Tyr Thr Ser Ala
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 Tyr Trp Cys Gly Gln Ala Leu Val Asp Val Ser Phe Phe Ile Leu Ile
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 Tyr Ala Ala Ser Leu Val Phe Phe Ile Tyr Met Ile Ser Phe Ile Phe
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 Arg Lys Arg Arg Lys Asn Ser Gly Leu Trp Ser Phe Tyr Phe Phe Phe
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 Ala Ser Thr Ile Met Phe Ser Ile Thr Leu Ile Asn His Phe Asp Leu
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 Ser Ile Leu Ile Thr Thr Met Val Leu Val Pro Ser Tyr Thr Leu Leu
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 Gly Phe Lys Thr Phe Leu Glu Val Arg Asp Gln Glu His Tyr Arg Glu
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 Phe Pro Glu Ala Asn Phe Glu Leu Ser Ala Thr Asp Phe Leu Val Cys

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 Met Glu Leu Lys Cys Gly Lys Lys Arg Met Arg Lys Asp Pro Val Phe
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 Arg Ile Ser Pro Gln Ser Arg Asp Ala Lys Pro Asn Pro Glu Glu Pro
 1235 1240 1245
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 1250 1255 1260
 Ala Leu Thr Thr Ser Ile Leu Asp Glu Lys Pro Val Ile Ile Ala Ser
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 Cys Leu His Lys Glu Tyr Ala Gly Gln Lys Lys Ser Cys Phe Ser Lys
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 Glu Leu Lys Gly Cys Ser Ser Val Leu Gly His Leu Gly Tyr Cys Pro
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 Gln Glu Asn Val Leu Trp Pro Met Leu Thr Leu Arg Glu His Leu Glu
 1365 1370 1375
 Val Tyr Ala Ala Val Lys Gly Leu Arg Lys Ala Asp Ala Arg Leu Ala
 1380 1385 1390
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 1395 1400 1405
 Pro Val Gln Lys Leu Thr Ala Gly Ile Thr Arg Lys Leu Cys Phe Val
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 Leu Ser Leu Leu Gly Asn Ser Pro Val Leu Leu Leu Asp Glu Pro Ser
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 Thr Gly Ile Asp Pro Thr Gly Gln Gln Gln Met Trp Gln Ala Ile Gln
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 1460 1465 1470
 Leu Ala Glu Ala Glu Ala Leu Cys Asp Arg Val Ala Ile Met Val Ser
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 Gly Arg Leu Arg Cys Ile Gly Ser Ile Gln His Leu Lys Asn Lys Leu
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 Gly Lys Asp Tyr Ile Leu Glu Leu Lys Val Lys Glu Thr Ser Gln Val
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Gln Glu Arg Tyr Ser Ser Leu Leu Thr Tyr Lys Leu Pro Val Ala Asp
 1540 1545 1550
 Val Tyr Pro Leu Ser Gln Thr Phe His Lys Leu Glu Ala Val Lys His
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 Asn Phe Asn Leu Glu Glu Tyr Ser Leu Ser Gln Cys Thr Leu Glu Lys
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 Val Phe Leu Glu Leu Ser Lys Glu Gln Glu Val Gly Asn Phe Asp Glu
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Pro

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 35 40 45
 Ser Asn Leu His Gln Val His Asp Thr Pro Gln Met Ser Ser Met Asp
 50 55 60
 Leu Gly Arg Val Asp Ser Phe Asn Asp Thr Asn Tyr Val Ile Ala Phe
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 Ala Pro Glu Ser Lys Thr Thr Gln Glu Ile Met Asn Lys Val Ala Ser
 85 90 95
 Ala Pro Phe Leu Lys Gly Arg Thr Ile Met Gly Trp Pro Asp Glu Lys
 100 105 110
 Ser Met Asp Glu Leu Asp Leu Asn Tyr Ser Ile Asp Ala Val Arg Val
 115 120 125
 Ile Phe Thr Asp Thr Phe Ser Tyr His Leu Lys Phe Ser Trp Gly His
 130 135 140
 Arg Ile Pro Met Met Lys Glu His Arg Asp His Ser Ala His Cys Gln
 145 150 155 160
 Ala Val Asn Glu Lys Met Lys Cys Glu Gly Ser Glu Phe Trp Glu Lys
 165 170 175
 Gly Phe Val Ala Phe Gln Ala Ala Ile Asn Ala Ala Ile Ile Glu Ile
 180 185 190
 Ala Thr Asn His Ser Val Met Glu Gln Leu Met Ser Val Thr Gly Val

195					200					205					
His	Met	Lys	Ile	Leu	Pro	Phe	Val	Ala	Gln	Gly	Gly	Val	Ala	Thr	Asp
	210					215					220				
Phe	Phe	Ile	Phe	Phe	Cys	Ile	Ile	Ser	Phe	Ser	Thr	Phe	Ile	Tyr	Tyr
225					230					235					240
Val	Ser	Val	Asn	Val	Thr	Gln	Glu	Arg	Gln	Tyr	Ile	Thr	Ser	Leu	Met
				245					250					255	
Thr	Met	Met	Gly	Leu	Arg	Glu	Ser	Ala	Phe	Trp	Leu	Ser	Trp	Gly	Leu
			260					265					270		
Met	Tyr	Ala	Gly	Phe	Ile	Leu	Ile	Met	Ala	Thr	Leu	Met	Ala	Leu	Ile
		275					280					285			
Val	Lys	Ser	Ala	Gln	Ile	Val	Val	Leu	Thr	Gly	Phe	Val	Met	Val	Phe
	290					295					300				
Thr	Leu	Phe	Leu	Leu	Tyr	Gly	Leu	Ser	Leu	Ile	Thr	Leu	Ala	Phe	Leu
305					310					315					320
Met	Ser	Val	Leu	Ile	Lys	Lys	Pro	Phe	Leu	Thr	Gly	Leu	Val	Val	Phe
				325					330					335	
Leu	Leu	Ile	Val	Phe	Trp	Gly	Ile	Leu	Gly	Phe	Pro	Ala	Leu	Tyr	Thr
			340					345					350		
His	Leu	Pro	Ala	Phe	Leu	Glu	Trp	Thr	Leu	Cys	Leu	Leu	Ser	Pro	Phe
		355					360					365			
Ala	Phe	Thr	Val	Gly	Met	Ala	Gln	Leu	Ile	His	Leu	Asp	Tyr	Asp	Val
	370					375					380				
Asn	Ser	Asn	Ala	His	Leu	Asp	Ser	Ser	Gln	Asn	Pro	Tyr	Leu	Ile	Ile
385					390					395					400
Ala	Thr	Leu	Phe	Met	Leu	Val	Phe	Asp	Thr	Leu	Leu	Tyr	Leu	Val	Leu
				405					410					415	
Thr	Leu	Tyr	Phe	Asp	Lys	Ile	Leu	Pro	Ala	Glu	Tyr	Gly	His	Arg	Cys
			420					425					430		
Ser	Pro	Leu	Phe	Phe	Leu	Lys	Ser	Cys	Phe	Trp	Phe	Gln	His	Gly	Arg
		435					440					445			
Ala	Asn	His	Val	Val	Leu	Glu	Asn	Glu	Thr	Asp	Ser	Asp	Pro	Thr	Pro
	450					455					460				
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Ile	Arg	Ile	Lys	Asn	Leu	Lys	Lys	Glu	Tyr	Ala	Gly	Lys	Cys	Glu	Arg
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Val	Glu	Ala	Leu	Lys	Gly	Val	Val	Phe	Asp	Ile	Tyr	Glu	Gly	Gln	Ile
			500					505					510		
Thr	Ala	Leu	Leu	Gly	His	Ser	Gly	Ala	Gly	Lys	Thr	Thr	Leu	Leu	Asn
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Asn His Thr Leu Ser Arg Met Ala Asp Ile Glu Asn Ile Ser Lys Phe
 545 550 555 560
 Thr Gly Phe Cys Pro Gln Ser Asn Val Gln Phe Gly Phe Leu Thr Val
 565 570 575
 Lys Glu Asn Leu Arg Leu Phe Ala Lys Ile Lys Gly Ile Leu Pro His
 580 585 590
 Glu Val Glu Lys Glu Val Gln Arg Val Val Gln Glu Leu Glu Met Glu
 595 600 605
 Asn Ile Gln Asp Ile Leu Ala Gln Asn Leu Ser Gly Gly Gln Asn Arg
 610 615 620
 Lys Leu Thr Phe Gly Ile Ala Ile Leu Gly Asp Pro Gln Val Leu Leu
 625 630 635 640
 Leu Asp Glu Pro Thr Ala Gly Leu Asp Pro Leu Ser Arg His Arg Ile
 645 650 655
 Trp Asn Leu Leu Lys Glu Gly Lys Ser Asp Arg Val Ile Leu Phe Ser
 660 665 670
 Thr Gln Phe Ile Asp Glu Ala Asp Ile Leu Ala Asp Arg Lys Val Phe
 675 680 685
 Ile Ser Asn Gly Lys Leu Lys Cys Ala Gly Ser Ser Leu Phe Leu Lys
 690 695 700
 Lys Lys Trp Gly Ile Gly Tyr His Leu Ser Leu His Leu Asn Glu Arg
 705 710 715 720
 Cys Asp Pro Glu Ser Ile Thr Ser Leu Val Lys Gln His Ile Ser Asp
 725 730 735
 Ala Lys Leu Thr Ala Gln Ser Glu Glu Lys Leu Val Tyr Ile Leu Pro
 740 745 750
 Leu Glu Arg Thr Asn Lys Phe Pro Glu Leu Tyr Arg Asp Leu Asp Arg
 755 760 765
 Cys Ser Asn Gln Gly Ile Glu Asp Tyr Gly Val Ser Ile Thr Thr Leu
 770 775 780
 Asn Glu Val Phe Leu Lys Leu Glu Gly Lys Ser Thr Ile Asp Glu Ser
 785 790 795 800
 Asp Ile Gly Ile Trp Gly Gln Leu Gln Thr Asp Gly Ala Lys Asp Ile
 805 810 815
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 820 825 830
 Arg Lys Thr Ile Ser Gly Val Ala Leu Trp Arg Gln Gln Val Cys Ala
 835 840 845
 Ile Ala Lys Val Arg Phe Leu Lys Leu Lys Lys Glu Arg Lys Ser Leu
 850 855 860
 Trp Thr Ile Leu Leu Leu Phe Gly Ile Ser Phe Ile Pro Gln Leu Leu
 865 870 875 880

Glu His Leu Phe Tyr Glu Ser Tyr Gln Lys Ser Tyr Pro Trp Glu Leu
 885 890 895
 Ser Pro Asn Thr Tyr Phe Leu Ser Pro Gly Gln Gln Pro Gln Asp Pro
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 Leu Thr His Leu Leu Val Ile Asn Lys Thr Gly Ser Thr Ile Asp Asn
 915 920 925
 Phe Leu His Ser Leu Arg Arg Gln Asn Ile Ala Ile Glu Val Asp Ala
 930 935 940
 Phe Gly Thr Arg Asn Gly Thr Asp Asp Pro Ser Tyr Asn Gly Ala Ile
 945 950 955 960
 Ile Val Ser Gly Asp Glu Lys Asp His Arg Phe Ser Ile Ala Cys Asn
 965 970 975
 Thr Lys Arg Leu Asn Cys Phe Pro Val Leu Leu Asp Val Ile Ser Asn
 980 985 990
 Gly Leu Leu Gly Ile Phe Asn Ser Ser Glu His Ile Gln Thr Asp Arg
 995 1000 1005
 Ser Thr Phe Phe Glu Glu His Met Asp Tyr Glu Tyr Gly Tyr Arg Ser
 1010 1015 1020
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 Ala Met Ser Ser Ile Gly Asp Tyr Lys Lys Lys Ala His Ser Gln Leu
 1045 1050 1055
 Arg Ile Ser Gly Leu Tyr Pro Ser Ala Tyr Trp Phe Gly Gln Ala Leu
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 Val Asp Val Ser Leu Tyr Phe Leu Ile Leu Leu Leu Met Gln Ile Met
 1075 1080 1085
 Asp Tyr Ile Phe Ser Pro Glu Glu Ile Ile Phe Ile Ile Gln Asn Leu
 1090 1095 1100
 Leu Ile Gln Ile Leu Cys Ser Ile Gly Tyr Val Ser Ser Leu Val Phe
 1105 1110 1115 1120
 Leu Thr Tyr Val Ile Ser Phe Ile Phe Arg Asn Gly Arg Lys Asn Ser
 1125 1130 1135
 Gly Ile Trp Ser Phe Phe Phe Leu Ile Val Val Ile Phe Ser Ile Val
 1140 1145 1150
 Ala Thr Asp Leu Asn Glu Tyr Gly Phe Leu Gly Leu Phe Phe Gly Thr
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 Met Leu Ile Pro Pro Phe Thr Leu Ile Gly Ser Leu Phe Ile Phe Ser
 1170 1175 1180
 Glu Ile Ser Pro Asp Ser Met Asp Tyr Leu Gly Ala Ser Glu Ser Glu
 1185 1190 1195 1200
 Ile Val Tyr Leu Ala Leu Leu Ile Pro Tyr Leu His Phe Leu Ile Phe
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 Leu Phe Ile Leu Arg Cys Leu Glu Met Asn Cys Arg Lys Lys Leu Met

1220	1225	1230
Arg Lys Asp Pro Val Phe Arg Ile Ser Pro Arg Ser Asn Ala Ile Phe 1235 1240 1245		
Pro Asn Pro Glu Glu Pro Glu Gly Glu Glu Glu Asp Ile Gln Met Glu 1250 1255 1260		
Arg Met Arg Thr Val Asn Ala Met Ala Val Arg Asp Phe Asp Glu Thr 1265 1270 1275 1280		
Pro Val Ile Ile Ala Ser Cys Leu Arg Lys Glu Tyr Ala Gly Lys Lys 1285 1290 1295		
Lys Asn Cys Phe Ser Lys Arg Lys Lys Thr Ile Ala Thr Arg Asn Val 1300 1305 1310		
Ser Phe Cys Val Lys Lys Gly Glu Val Ile Gly Leu Leu Gly His Asn 1315 1320 1325		
Gly Ala Gly Lys Ser Thr Thr Ile Lys Met Ile Thr Gly Asp Thr Lys 1330 1335 1340		
Pro Thr Ala Gly Gln Val Ile Leu Lys Gly Ser Gly Gly Gly Glu Pro 1345 1350 1355 1360		
Leu Gly Phe Leu Gly Tyr Cys Pro Gln Glu Asn Ala Leu Trp Pro Asn 1365 1370 1375		
Leu Thr Val Arg Gln His Leu Glu Val Tyr Ala Ala Val Lys Gly Leu 1380 1385 1390		
Arg Lys Gly Asp Ala Met Ile Ala Ile Thr Arg Leu Val Asp Ala Leu 1395 1400 1405		
Lys Leu Gln Asp Gln Leu Lys Ala Pro Val Lys Thr Leu Ser Glu Gly 1410 1415 1420		
Ile Lys Arg Lys Leu Arg Phe Val Leu Ser Ile Leu Gly Asn Pro Ser 1425 1430 1435 1440		
Val Val Leu Leu Asp Glu Pro Ser Thr Gly Met Asp Pro Glu Gly Gln 1445 1450 1455		
Gln Gln Met Trp Gln Val Ile Arg Ala Thr Phe Arg Asn Thr Glu Arg 1460 1465 1470		
Gly Ala Leu Leu Thr Thr His Tyr Met Ala Glu Ala Glu Ala Val Cys 1475 1480 1485		
Asp Arg Val Ala Ile Met Val Ser Gly Arg Leu Arg Cys Ile Gly Ser 1490 1495 1500		
Ile Gln His Leu Lys Ser Lys Phe Gly Lys Asp Tyr Leu Leu Glu Met 1505 1510 1515 1520		
Lys Leu Lys Asn Leu Ala Gln Met Glu Pro Leu His Ala Glu Ile Leu 1525 1530 1535		
Arg Leu Phe Pro Gln Ala Ala Gln Gln Glu Arg Phe Ser Ser Leu Met 1540 1545 1550		
Val Tyr Lys Leu Pro Val Glu Asp Val Arg Pro Leu Ser Gln Ala Phe 1555 1560 1565		

Phe Lys Leu Glu Ile Val Lys Gln Ser Phe Asp Leu Glu Glu Tyr Ser
 1570 1575 1580
 Leu Ser Gln Ser Thr Leu Glu Gln Val Phe Leu Glu Leu Ser Lys Glu
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 Lys Leu Leu Leu Gln Glu Glu Pro
 1620

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 <213> Homo sapiens

<220>
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 20 25 30
 His Glu Met Val Gly Val Ile Phe Ser Asp Thr Phe Ser Tyr Arg Leu
 35 40 45
 Lys Phe Asn Trp Gly Tyr Arg Ile Pro Val Ile Lys Glu His Ser Glu
 50 55 60
 Tyr Thr Glu His Cys Trp Ala Met His Gly Glu Ile Phe Cys Tyr Leu
 65 70 75 80
 Ala Lys Tyr Trp Leu Lys Gly Phe Val Ala Phe Gln Ala Ala Ile Asn
 85 90 95
 Ala Ala Ile Ile Glu Val Thr Thr Asn His Ser Val Met Glu Glu Leu
 100 105 110
 Thr Ser Val Ile Gly Ile Asn Met Lys Ile Pro Pro Phe Ile Ser Lys
 115 120 125
 Gly Glu Ile Met Asn Glu Trp Phe His Phe Thr Cys Leu Val Ser Phe
 130 135 140
 Ser Ser Phe Ile Tyr Phe Ala Ser Leu Asn Val Ala Arg Glu Arg Gly
 145 150 155 160
 Lys Phe Lys Lys Leu Met Thr Val Met Gly Leu Arg Glu Ser Ala Phe
 165 170 175
 Trp Leu Ser Trp Xaa Leu Thr Tyr Ile Cys Phe Ile Phe Ile Met Ser
 180 185 190
 Ile Phe Met Ala Leu Val Ile Thr Ser Ile Ser Ile Val Phe His Thr
 195 200 205

Gly Phe Met Val Ile Phe Thr Leu Tyr Ser Leu Tyr Gly Leu Ser Leu
 210 215 220
 Ile Ala Leu Ala Phe Leu Met Ser Val Leu Ile Arg Lys Pro Met Leu
 225 230 235 240
 Ala Gly Leu Ala Gly Phe Leu Phe Thr Val Phe Trp Gly Cys Leu Gly
 245 250 255
 Phe Thr Val Leu Tyr Arg Gln Leu Pro Leu Ser Leu Gly Trp Val Leu
 260 265 270
 Ser Leu Leu Ser Pro Phe Ala Phe Thr Ala Gly Met Ala Gln Val Thr
 275 280 285
 His Leu Asp Asn Tyr Leu Ser Gly Val Ile Phe Pro Asp Pro Ser Gly
 290 295 300
 Asp Ser Tyr Lys Met Ile Ala Thr Phe Phe Ile Leu Ala Phe Asp Thr
 305 310 315 320
 Leu Phe Tyr Leu Ile Phe Thr Leu Tyr Phe Glu Arg Val Leu Pro Asp
 325 330 335
 Lys Asp Gly His Gly Asp Ser Pro Leu Phe Phe Leu Lys Ser Ser Phe
 340 345 350
 Trp Ser Lys His Gln Asn Thr His His Glu Ile Phe Glu Asn Glu Ile
 355 360 365
 Asn Pro Glu His Ser Ser Asp Asp Ser Phe Glu Pro Val Ser Pro Glu
 370 375 380
 Phe His Gly Lys Glu Ala Ile Arg Ile Arg Asn Val Ile Lys Glu Tyr
 385 390 395 400
 Asn Gly Lys Thr Gly Lys Val Glu Ala Leu Gln Gly Ile Phe Phe Asp
 405 410 415
 Ile Tyr Glu Gly Gln Ile Thr Ala Ile Leu Gly His Asn Gly Ala Gly
 420 425 430
 Lys Ser Thr Leu Leu Asn Ile Leu Ser Gly Leu Ser Val Ser Thr Glu
 435 440 445
 Gly Ser Ala Thr Ile Tyr Asn Thr Gln Leu Ser Glu Ile Thr Asp Met
 450 455 460
 Glu Glu Ile Arg Lys Asn Ile Gly Phe Cys Pro Gln Phe Asn Phe Gln
 465 470 475 480
 Phe Asp Phe Leu Thr Val Arg Glu Asn Leu Arg Val Phe Ala Lys Ile
 485 490 495
 Lys Gly Ile Gln Pro Lys Glu Val Glu Gln Glu Val Lys Arg Ile Ile
 500 505 510
 Met Glu Leu Asp Met Gln Ser Ile Gln Asp Ile Ile Ala Lys Lys Leu
 515 520 525
 Ser Gly Gly Gln Lys Arg Lys Leu Thr Leu Gly Ile Ala Ile Leu Gly
 530 535 540

Asp Pro Gln Val Leu Leu Leu Asp Glu Pro Thr Ala Gly Leu Asp Pro
 545 550 555 560
 Phe Ser Arg His Arg Val Trp Ser Leu Leu Lys Glu His Lys Val Asp
 565 570 575
 Arg Leu Ile Leu Phe Ser Thr Gln Phe Met Asp Glu Ala Asp Ile Leu
 580 585 590
 Ala Asp Arg Lys Val Phe Leu Ser Asn Gly Lys Leu Lys Cys Ala Gly
 595 600 605
 Ser Ser Leu Phe Leu Lys Arg Lys Trp Gly Ile Gly Tyr His Leu Ser
 610 615 620
 Leu His Arg Asn Glu Met Cys Asp Thr Glu Lys Ile Thr Ser Leu Ile
 625 630 635
 Lys Gln His Ile Pro Asp Ala Lys Leu Thr Thr Glu Ser Glu Glu Lys
 645 650 655
 Leu Val Tyr Ser Leu Pro Leu Glu Lys Thr Asn Lys Phe Pro Asp Leu
 660 665 670
 Tyr Ser Asp Leu Asp Lys Cys Ser Asp Gln Gly Ile Arg Asn Tyr Ala
 675 680 685
 Val Ser Val Thr Ser Leu Asn Glu Val Phe Leu Asn Leu Glu Gly Lys
 690 695 700
 Ser Ala Ile Asp Glu Pro Asp Phe Asp Ile Gly Lys Gln Glu Lys Ile
 705 710 715 720
 His Val Thr Arg Asn Thr Gly Asp Glu Ser Glu Met Glu Gln Val Leu
 725 730 735
 Cys Ser Leu Pro Glu Thr Arg Lys Ala Val Ser Ser Ala Ala Leu Trp
 740 745 750
 Arg Arg Gln Ile Tyr Ala Val Ala Thr Leu Arg Phe Leu Lys Leu Arg
 755 760 765
 Arg Glu Arg Arg Ala Leu Leu Cys Leu Leu Leu Val Leu Gly Ile Ala
 770 775 780
 Phe Ile Pro Ile Ile Leu Glu Lys Ile Met Tyr Lys Val Thr Arg Glu
 785 790 795 800
 Thr His Cys Trp Glu Phe Ser Pro Ser Met Tyr Phe Leu Ser Leu Glu
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 820 825 830
 Gly Ser Asn Ile Glu Asp Leu Val His Ser Leu Lys Cys Gln Asp Ile
 835 840 845
 Val Leu Glu Ile Asp Asp Phe Arg Asn Arg Asn Gly Ser Asp Asp Pro
 850 855 860
 Ser Tyr Asn Gly Ala Ile Ile Val Ser Gly Asp Gln Lys Asp Tyr Arg
 865 870 875 880
 Phe Ser Val Ala Cys Asn Thr Lys Lys Leu Asn Cys Phe Pro Val Leu

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Leu	Ile	Gln	Thr	Glu	Ser	Thr	Ser	Phe	Ser	Arg	Asp	Asp	Ile	Val	Leu				
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	930					935					940								
Cys	Val	Ser	Pro	Phe	Ile	Gly	Met	Ser	Ser	Ile	Ser	Asp	Tyr	Lys	Lys				
945					950					955					960				
Asn	Val	Gln	Ser	Gln	Leu	Trp	Ile	Ser	Gly	Leu	Trp	Pro	Ser	Ala	Tyr				
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Trp	Cys	Gly	Gln	Ala	Leu	Val	Asp	Ile	Pro	Leu	Tyr	Phe	Leu	Ile	Leu				
			980					985					990						
Phe	Ser	Ile	His	Leu	Ile	Tyr	Tyr	Phe	Ile	Phe	Leu	Gly	Phe	Gln	Leu				
		995				1000						1005							
Ser	Trp	Glu	Leu	Met	Phe	Val	Leu	Val	Val	Cys	Ile	Ile	Gly	Cys	Ala				
	1010				1015					1020									
Val	Ser	Leu	Ile	Phe	Leu	Thr	Tyr	Val	Leu	Ser	Phe	Ile	Phe	Arg	Lys				
1025				1030					1035					1040					
Trp	Arg	Lys	Asn	Asn	Gly	Phe	Trp	Ser	Phe	Gly	Phe	Phe	Ile	Ile	Leu				
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Ile	Cys	Val	Ser	Thr	Ile	Met	Val	Ser	Thr	Gln	Tyr	Glu	Lys	Leu	Asn				
		1060					1065					1070							
Leu	Ile	Leu	Cys	Met	Ile	Phe	Ile	Pro	Ser	Phe	Thr	Leu	Leu	Gly	Tyr				
	1075				1080						1085								
Val	Met	Leu	Leu	Ile	Gln	Leu	Asp	Phe	Met	Arg	Asn	Leu	Asp	Ser	Leu				
	1090				1095						1100								
Asp	Asn	Arg	Ile	Asn	Glu	Val	Asn	Lys	Thr	Ile	Leu	Leu	Thr	Thr	Leu				
1105				1110					1115					1120					
Ile	Pro	Tyr	Leu	Gln	Ser	Val	Ile	Phe	Leu	Phe	Val	Ile	Arg	Cys	Leu				
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Glu	Met	Lys	Tyr	Gly	Asn	Glu	Ile	Met	Asn	Lys	Asp	Pro	Val	Phe	Arg				
		1140					1145					1150							
Ile	Ser	Pro	Arg	Ser	Arg	Glu	Thr	His	Pro	Asn	Pro	Glu	Glu	Pro	Glu				
	1155					1160						1165							
Glu	Glu	Asp	Glu	Asp	Val	Gln	Ala	Glu	Arg	Val	Gln	Ala	Ala	Asn	Ala				
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Leu	Thr	Ala	Pro	Asn	Leu	Glu	Glu	Glu	Pro	Val	Ile	Thr	Ala	Ser	Cys				
1185				1190					1195					1200					
Leu	His	Lys	Glu	Tyr	Tyr	Glu	Thr	Lys	Lys	Ser	Cys	Phe	Ser	Thr	Arg				
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Lys	Lys	Lys	Ile	Ala	Ile	Arg	Asn	Val	Ser	Phe	Cys	Val	Lys	Lys	Gly				
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 Ile Lys Met Ile Thr Gly Cys Thr Lys Pro Thr Ala Gly Val Val Val
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 Leu Gln Gly Ser Arg Ala Ser Val Arg Gln Gln His Asp Asn Ser Leu
 1265 1270 1275 1280
 Lys Phe Leu Gly Tyr Cys Pro Gln Glu Asn Ser Leu Trp Pro Lys Leu
 1285 1290 1295
 Thr Met Lys Glu His Leu Glu Leu Tyr Ala Ala Val Lys Gly Leu Gly
 1300 1305 1310
 Lys Glu Asp Ala Ala Leu Ser Ile Ser Arg Leu Val Glu Ala Leu Lys
 1315 1320 1325
 Leu Gln Glu Gln Leu Lys Ala Pro Val Lys Thr Leu Ser Glu Gly Ile
 1330 1335 1340
 Lys Arg Lys Leu Cys Phe Val Leu Ser Ile Leu Gly Asn Pro Ser Val
 1345 1350 1355 1360
 Val Leu Leu Asp Glu Pro Phe Thr Gly Met Asp Pro Glu Gly Gln Gln
 1365 1370 1375
 Gln Met Trp Gln Ile Leu Gln Ala Thr Val Lys Asn Lys Glu Arg Gly
 1380 1385 1390
 Thr Leu Leu Thr Thr His Tyr Met Ser Glu Ala Glu Ala Val Cys Asp
 1395 1400 1405
 Arg Met Ala Met Met Val Ser Gly Thr Leu Arg Cys Ile Gly Ser Ile
 1410 1415 1420
 Gln His Leu Lys Asn Lys Phe Gly Arg Asp Tyr Leu Leu Glu Ile Lys
 1425 1430 1435 1440
 Met Lys Glu Pro Thr Gln Val Glu Ala Leu His Thr Glu Ile Leu Lys
 1445 1450 1455
 Leu Phe Pro Gln Ala Ala Trp Gln Glu Arg Tyr Ser Ser Leu Met Ala
 1460 1465 1470
 Tyr Lys Leu Pro Val Glu Asp Val His Pro Leu Ser Arg Ala Phe Phe
 1475 1480 1485
 Lys Leu Glu Ala Met Lys Gln Thr Phe Asn Leu Glu Glu Tyr Ser Leu
 1490 1495 1500
 Ser Gln Ala Thr Leu Glu Gln Val Phe Leu Glu Leu Cys Lys Glu Gln
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<213> Homo sapiens

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<211> 141
<212> DNA
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atgaaaagag agagcttatt g 141

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<211> 205
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tcttctttta tggttgtgta tacaccaata tctaatttaa cccagcagat aatgaataaa 180
acagcacttg ctctctttt gaaag 205

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<211> 159
<212> DNA
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agggatataa cagtccactt tggaagaag atttctcag 159

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<212> DNA
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<210> 14
<211> 227
<212> DNA
<213> Homo sapiens

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cttcatttct cccacttgt atattttata tcaactcaatg taacaaaaga gagaaaaaag 180
tctaagaatt tgatgaaaat gatgggtctc caagattcag cattctg 227

<210> 15
<211> 142
<212> DNA
<213> Homo sapiens

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ttttttatat ggcttatctt tg 142

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<211> 186
<212> DNA
<213> Homo sapiens

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attcag 186

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<211> 148
<212> DNA
<213> Homo sapiens

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tatacaatga tagcaacttt ttctatgttg cttttggatg gtctcatcta cttgctattg 120
gcattatact ttgacaaaat ttaccct 148

<210> 18
<211> 169
<212> DNA
<213> Homo sapiens

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accaaaggac taatgctaag gttattgaga aagaaatcga tgctgagcat ccctctgatg 120
attattttga accagtagct cctgaattcc aaggaaaaga agccatcag 169

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<211> 59
<212> DNA
<213> Homo sapiens

<400> 19
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<212> DNA
<213> Homo sapiens

<400> 20
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<210> 21
 <211> 176
 <212> DNA
 <213> Homo sapiens

<400> 21
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 agataactgg cgtctgtcct caattcaatg ttcaatttga catactcacc gtgaaggaaa 120
 acctcagcct gtttgctaaa ataaaagga ttcattctaaa ggaagtggaa caagag 176

<210> 22
 <211> 120
 <212> DNA
 <213> Homo sapiens

<400> 22
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 ttaagtgaag gacagaaaag aaagctgact tttgggatta ccatttttagg agatcctcaa 120

<210> 23
 <211> 139
 <212> DNA
 <213> Homo sapiens

<400> 23
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 gaggctgaca tcctggctg 139

<210> 24
 <211> 91
 <212> DNA
 <213> Homo sapiens

<400> 24
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 aaagaagggt gggctcttga taccaccta g 91

<210> 25
 <211> 140
 <212> DNA
 <213> Homo sapiens

<400> 25
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 aaggacaaat acatttccag 140

<210> 26
 <211> 117
 <212> DNA
 <213> Homo sapiens

<400> 26
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 tgtcaactct aaatgaagtc tttatgaaac tggaaggaca gtcaactatc gaacaag 117

<210> 27
<211> 184
<212> DNA
<213> Homo sapiens

<400> 27
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tctttgccat ggcacggctc cgtttcttaa agttaaaacg tcaaactaaa gtgttattga 180
ccct 184

<210> 28
<211> 167
<212> DNA
<213> Homo sapiens

<400> 28
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gttaaatgaa aagatcgatt gggaatttaa aaacgaattg tattttctct ctcctggaca 120
acttccccag gaaccccgta ccagcctggt gatcatcaat aacacag 167

<210> 29
<211> 134
<212> DNA
<213> Homo sapiens

<400> 29
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ctggtaaaca aaag 134

<210> 30
<211> 138
<212> DNA
<213> Homo sapiens

<400> 30
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aatattatca gcaatgggct acttcaaagt tttaatcaca cacaacatat tcgaattgag 120
tcaagcccat ttcctctt 138

<210> 31
<211> 108
<212> DNA
<213> Homo sapiens

<400> 31
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tgtagcattt ctccttatat caccatgggc agcatcagtg attacaag 108

<210> 32
<211> 174
<212> DNA
<213> Homo sapiens

<400> 32
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caggcactag tggacgtcag cttcttcatt ttaattctcc ttttaatgta ttttaatttc 120

tacatagaaa acatgcagta ctttcttatt acaagccaaa ttgtgtttgc ttg 174

<210> 33
<211> 114
<212> DNA
<213> Homo sapiens

<400> 33
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atttttcgca aaaggagaaa aaacagtggc ctttgggtcat ttacttctt tttt 114

<210> 34
<211> 120
<212> DNA
<213> Homo sapiens

<400> 34
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accaccatgg tattgggttc ttcataatcc ttgcttggat ttaaaacttt tttggaagt 120

<210> 35
<211> 78
<212> DNA
<213> Homo sapiens

<400> 35
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tttctagtct gtttcata 78

<210> 36
<211> 92
<212> DNA
<213> Homo sapiens

<400> 36
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aagaaaagaa tgcgaaaaga tctgttttc ag 92

<210> 37
<211> 121
<212> DNA
<213> Homo sapiens

<400> 37
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agatattcaa acagaaagaa taagaacagc cactgtcttg accacttcaa tcttagatga 120
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<210> 38
<211> 118
<212> DNA
<213> Homo sapiens

<400> 38
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<210> 39
<211> 92
<212> DNA
<213> Homo sapiens

<400> 39
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tatctgggat cacaaagcca actgctggag ag 92

<210> 40
<211> 155
<212> DNA
<213> Homo sapiens

<400> 40
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gtgctgtggc ccatgctgac gttgagggaa cacctggagg tgtatgctgc cgtcaagggg 120
ctcaggaaag cggacgcgag gctcgccatc gcaag 155

<210> 41
<211> 76
<212> DNA
<213> Homo sapiens

<400> 41
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aggaatcacg agaaag 76

<210> 42
<211> 95
<212> DNA
<213> Homo sapiens

<400> 42
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acgggcatag accccacagg gcagcagcaa atgtg 95

<210> 43
<211> 120
<212> DNA
<213> Homo sapiens

<400> 43
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cctggctgag gcggaagcct tgtgtgaccg tgtggccatc atggtgtctg gaaggcttag 120

<210> 44
<211> 141
<212> DNA
<213> Homo sapiens

<400> 44
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aaaagtgaag gaaacgtctc aagtgacttt ggtccacact gagattctga agcttttccc 120
acaggctgca gggcaggaaa g 141

<210> 45
<211> 80

<212> DNA
<213> Homo sapiens

<400> 45
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ctttcacaaa ttagaagcag 80

<210> 46
<211> 56
<212> DNA
<213> Homo sapiens

<400> 46
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<210> 47
<211> 369
<212> DNA
<213> Homo sapiens

<400> 47
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acaatgagat ggaaactcct ccctcattca gatgaacctt aaaacctcaa acctagtaat 120
tttttggtga tctcctataa acttatgttt tatgtaataa ttaatagtat gtttaatttt 180
aaagatcatt taaaattaac atcagggtata ttttgtaaatt ttagttaaca aatacataaa 240
ttttaaaatt attcttcctc tcaaaccatag ggggtgatagc aaacctgtga taaaggcaat 300
acaaaatatt agtaaagtca cccaaagagt caggcactgg gtattgtgga aataaaaacta 360
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<210> 48
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<212> DNA
<213> Homo sapiens

<400> 48
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tctcctccag aacatgcaga gacccatgga tgaactgtgt ttctagattt ttctccagc 120
tttcttgaga 130

<210> 49
<211> 109
<212> DNA
<213> Homo sapiens

<400> 49
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ctgcaagaac tgtctcaaaa aatggagaat gaaaagacag accttggtg 109

<210> 50
<211> 208
<212> DNA
<213> Homo sapiens

<400> 50
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caagttcatg acactcctca aatgtcttca atggatctgg gacgtgtaga tagttttaat 120
gatactaatt atgttattgc atttgcacct gaatccaaaa ctaccaaga gataatgaac 180
aaagtggctt cagccccatt cctaaaag 208

<210> 51
<211> 165
<212> DNA
<213> Homo sapiens

<400> 51
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caatagacgc agtgagagtc atctttactg ataccttctc ctaccatttg aagttttctt 120
ggggacatag aatcccatg atgaaagagc acagagacca ttcag 165

<210> 52
<211> 104
<212> DNA
<213> Homo sapiens

<400> 52
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gctttgtagc ttttcaagct gccattaatg ctgctatcat agaa 104

<210> 53
<211> 227
<212> DNA
<213> Homo sapiens

<400> 53
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atattacctt ttgttgccca aggaggagtt gcaactgatt ttttcatttt cttttgcatt 120
atctcttttt ctacatttat atactatgta tcagtcaatg ttacacaaga aagacaatac 180
attacgtcat tgatgacaat gatgggactc cgagagtcag cattctg 227

<210> 54
<211> 142
<212> DNA
<213> Homo sapiens

<400> 54
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tattgtaaaa tctgcacaaa ttgtcgtcct gactgggttt gtgatgggtc tcaccctctt 120
tctcctctat ggcctgtctt tg 142

<210> 55
<211> 186
<212> DNA
<213> Homo sapiens

<400> 55
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gtgtttctcc ttattgtctt ttgggggatc ctgggattcc cagcattgta tacacatctt 120
cctgcatttt tggaatggac tttgtgtctt cttagccccct ttgccttcac tgttgggatg 180
gccag 186

<210> 56
<211> 148
<212> DNA
<213> Homo sapiens

<400> 56
cttatacatt tggactatga tgtgaattct aatgccact tggattcttc acaaaatcca 60

tacctcataa tagctactct tttcatgttg gtttttgaca cccttctgta tttggtattg 120
acattatatt ttgacaaaat tttgcccg 148

<210> 57
<211> 169
<212> DNA
<213> Homo sapiens

<400> 57
ctgaatatgg acatcgatgt tctcccttgt ttttcctgaa atcctgtttt tggtttcaac 60
acggaagggc taatcatgtg gtccttgaga atgaaacaga ttctgacct acacctaag 120
actgttttga accagtgtct ccagaattct gtgggaagga agccatcag 169

<210> 58
<211> 59
<212> DNA
<213> Homo sapiens

<400> 58
aatcaaaaat cttaaaaaag aatatgcagg gaagtgtgag agagtagaag ctttgaaag 59

<210> 59
<211> 111
<212> DNA
<213> Homo sapiens

<400> 59
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gaaaaactac cctgttaaag atacttagtg ggttgctcagt tccaacatca g 111

<210> 60
<211> 176
<212> DNA
<213> Homo sapiens

<400> 60
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agttcactgg attttgtcca caatccaatg tgcaatttgg atttctcact gtgaaagaaa 120
acctcaggct gtttgctaaa ataaaagggg ttttgccaca tgaagtggag aaagag 176

<210> 61
<211> 120
<212> DNA
<213> Homo sapiens

<400> 61
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ttaagtgggt gacaaaatag gaaactaact tttgggattg ccattttagg agatcctcaa 120

<210> 62
<211> 139
<212> DNA
<213> Homo sapiens

<400> 62
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aatctcctga aagaggggaa atcagacaga gtaattctct tcagcaccga gtttatagat 120
gaggctgaca ttctggcgg 139

<210> 63
<211> 91
<212> DNA
<213> Homo sapiens

<400> 63
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agaagaaatg gggcataggc taccatttaa g 91

<210> 64
<211> 140
<212> DNA
<213> Homo sapiens

<400> 64
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ctctgatgcc aaattgacag cacaaagtga agaaaaactt gtatatattt tgcctttgga 120
aaggacaaac aaatttccag 140

<210> 65
<211> 120
<212> DNA
<213> Homo sapiens

<400> 65
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taacaacttt gaatgaggtg tttctgaaat tagaaggaaa atcaactatt gatgaatcag 120

<210> 66
<211> 199
<212> DNA
<213> Homo sapiens

<400> 66
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agctggaaca agttttgtct tccttccacg aaacaaggaa aacaatcagt ggcgtggcgc 120
tctggaggca gcaggtctgt gcaatagcaa aagttcgctt cctaaagtta aagaaagaaa 180
gaaaaagcct gtggactat 199

<210> 67
<211> 167
<212> DNA
<213> Homo sapiens

<400> 67
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atatcagaaa agttaccctg gggaactgtc tccaaataca tacttcctct caccaggaca 120
acaaccacag gatcctctga cccatttact ggtcatcaat aagacag 167

<210> 68
<211> 134
<212> DNA
<213> Homo sapiens

<400> 68
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atgccttttg aactagaaat ggcacagatg acccatctta caatgggtgct atcattgtgt 120

caggtgatga aaag

134

<210> 69
<211> 138
<212> DNA
<213> Homo sapiens

<400> 69
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gatgtcatta gcaatggact acttgggaatt ttttaattcgt cagaacacat tcagactgac 120
agaagcacat tttttgaa 138

<210> 70
<211> 108
<212> DNA
<213> Homo sapiens

<400> 70
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gcctctttca ctccatacat tgcaatgagc agcattgggtg actacaaa 108

<210> 71
<211> 174
<212> DNA
<213> Homo sapiens

<400> 71
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caagcactgg tggatgtttc cctgtacttt ttgatcctcc tgctaatagca aataatggat 120
tatattttta gcccagagga gatttatattt ataattcaaa acctgttaat tcaa 174

<210> 72
<211> 114
<212> DNA
<213> Homo sapiens

<400> 72
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atctttcgca atgggagaaa aaatagtggc atttgggtcat ttttcttctt aatt 114

<210> 73
<211> 120
<212> DNA
<213> Homo sapiens

<400> 73
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<210> 74
<211> 69
<212> DNA
<213> Homo sapiens

<400> 74
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ctgctaata 69

<210> 75
<211> 92
<212> DNA
<213> Homo sapiens

<400> 75
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aagaaactaa tgagaaagga tcctgtgttc ag 92

<210> 76
<211> 121
<212> DNA
<213> Homo sapiens

<400> 76
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g 121

<210> 77
<211> 118
<212> DNA
<213> Homo sapiens

<400> 77
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ttttctaaaa ggaagaaaac aattgccaca agaatgtct ctttttgtgt taaaaaag 118

<210> 78
<211> 92
<212> DNA
<213> Homo sapiens

<400> 78
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<210> 79
<211> 161
<212> DNA
<213> Homo sapiens

<400> 79
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aaaggtctca ggaaagggga cgcaatgatc gccatcacac g 161

<210> 80
<211> 76
<212> DNA
<213> Homo sapiens

<400> 80
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gggaataaag cgaaag 76

<210> 81

<211> 95
<212> DNA
<213> Homo sapiens

<400> 81
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accgggatgg accccgaggg gcagcagcaa atgtg 95

<210> 82
<211> 120
<212> DNA
<213> Homo sapiens

<400> 82
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catggcagag gctgaggcgg tgtgtgaccg agtggccatc atggtgtcag gaaggctgag 120

<210> 83
<211> 141
<212> DNA
<213> Homo sapiens

<400> 83
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gaagctgaag aacctggcac aaatggagcc cctccatgca gagatcctga ggcttttccc 120
ccaggctgct cagcaggaaa g 141

<210> 84
<211> 80
<212> DNA
<213> Homo sapiens

<400> 84
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tttcttcaaa ttagagatatg 80

<210> 85
<211> 56
<212> DNA
<213> Homo sapiens

<400> 85
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<210> 86
<211> 1062
<212> DNA
<213> Homo sapiens

<400> 86
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tctgtatcat	tctactataa	agacacatgc	acacgtatgt	ttattgcagc	actgtttaca	720
atagcaaaga	cttggaacca	accaaataac	ccacaaatga	tagaccggat	aaagaaaacg	780
tgacacatat	acaccatgga	atactatgca	gccatagaaa	aggatgagtt	catattcttc	840
acagggacat	ggatgaagct	ggaaaccatc	atcctcagca	aactaacaca	ggaacagaaa	900
accaaacacc	gcatgtttct	actcataagt	gggaattgaa	caatgagaat	acatggacac	960
agggagggga	acaccacacc	ctggggcctg	ttggggggat	gggggctagg	ggagggatag	1020
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<210> 87
 <211> 287
 <212> DNA
 <213> Homo sapiens

<400> 87						
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ctcaaaactt	cattctaatt	gtgccctgag	tttgttaaaa	taccatactg	tattttttgtg	120
taacatgtaa	ataggcatta	atttttgaga	aatagaaatg	tttatcctta	atgtattttt	180
aatttgctaa	cattgatatt	ttattttctt	tcctgaaata	gcttattttc	taaaatgaaa	240
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<210> 88
 <211> 280
 <212> DNA
 <213> Homo sapiens

<400> 88						
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aactggacaa	ggagaaaaac	atagggaaaa	aaccaacaga	atttgttggc	atgttctaca	120
cacagaccat	ggctttttcag	aagccaagct	gaataaaaaac	agtttttaaaa	gaggcaacca	180
tttgtagagg	agtccttgaa	ggattcttca	ttgtttttctt	ggacaaaaag	agaccagtgg	240
atccaagtgc	ttcaaatact	tctctcttat	tttcttaact			280

<210> 89
 <211> 141
 <212> DNA
 <213> Homo sapiens

<400> 89						
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agataagcgt	gcgtcaacaa	attcaggctc	ttctgtacaa	gaattttctt	aaaaaatgga	120
gaataaaaag	agagttttatt	g				141

<210> 90
 <211> 205
 <212> DNA
 <213> Homo sapiens

<400> 90						
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gctaccggtt	ttcctgaaca	acctcctaaa	gtcctgggaa	gcgtggatca	gtttaatgac	120
tctggcctgg	tagtggcata	tacaccagtc	agtaacataa	cacaaaggat	aatgaataag	180
atggccttgg	cttcctttat	gaaag				205

<210> 91
 <211> 165
 <212> DNA
 <213> Homo sapiens

<400> 91
gaagaacagt cattgggaca ccagatgaag agaccatgga tatagaactt ccaaaaaaat 60
accatgaaat ggtgggagtt atatttagtg atactttctc atatcgctg aagtttaatt 120
gggatatag aatcccagtt ataaaggagc actctgaata cacag 165

<210> 92
<211> 104
<212> DNA
<213> Homo sapiens

<400> 92
aacactgttg ggccatgcat ggtgaaattt tttgttactt ggcaaagtac tggctaaaag 60
ggtttgtagc ttttcaagct gcaattaatg ctgcaattat agaa 104

<210> 93
<211> 227
<212> DNA
<213> Homo sapiens

<400> 93
gtcacaacaa atcattctgt aatggaggag ttgacatcag ttatttgaat aaatatgaag 60
ataccacctt tcatttctaa gggagaaatt atgaatgaat ggtttcattt tacttgctta 120
gtttctttct cttcttttat atactttgca tcattaaatg ttgcaaggga aagaggaaaa 180
tttaagaaac tgatgacagt aatgggtctc cgagagtcag cattctg 227

<210> 94
<211> 142
<212> DNA
<213> Homo sapiens

<220>
<221> unsure
<222> 11
<223> n=unknown, may be a or g or c or t

<400> 94
gctctccttg ngattgacat acatttgctt catcttcatt atgtccattt ttatggctct 60
ggtcataaca tcaatctcaa ttgtatttca tactggcttc atggtgatat tcacactcta 120
tagcttatat ggcctttctt tg 142

<210> 95
<211> 186
<212> DNA
<213> Homo sapiens

<400> 95
atagcatttg ctttcctcat gagtgtttta ataaggaaac ctatgctcgc tggtttggct 60
ggatttctct tcaactgtatt ttggggatgt ctgggattca ctgtgttata cagacaactt 120
cctttatctt tgggatgggt attaatgtct cttagccctt ttgccttcac tgctggaatg 180
gccag 186

<210> 96
<211> 148
<212> DNA
<213> Homo sapiens

<400> 96
gttacacacc tggataatta cttaagtggg gttatttttc ctgatccctc tggggattca 60

tacaaaatga tagccacttt tttcattttg gcatttgata ctcttttcta tttgatattc 120
acattatatt ttgagcgagt tttacctg 148

<210> 97
<211> 169
<212> DNA
<213> Homo sapiens

<400> 97
ataaagatgg ccatggggat tctccattat ttttccttaa gtcctcattt tgggtccaaac 60
atcaaaatac tcatcatgaa atcttttgaga atgaaataaa tcctgagcat tcctctgatg 120
attcctttga accggtgtct ccagaattcc atggaaaaga agccataag 169

<210> 98
<211> 59
<212> DNA
<213> Homo sapiens

<400> 98
aatcagaaat gttataaaaag aatataatgg aaagactgga aaagtagaag cattgcaag 59

<210> 99
<211> 111
<212> DNA
<213> Homo sapiens

<400> 99
gcataatatt tgacatatat gaaggacaga tcactgcaat acttgggcat aatggagctg 60
gtaaatcaac actgctaacc attcttagtg gattgtctgt ttctacagaa g 111

<210> 100
<211> 176
<212> DNA
<213> Homo sapiens

<400> 100
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agaatatttg attttgtcca cagttcaatt ttcaatttga cttcctcact gtgagagaaa 120
acctcagggg atttgctaaa ataaaagggg ttcagccaaa ggaagtggaa caagag 176

<210> 101
<211> 120
<212> DNA
<213> Homo sapiens

<400> 101
gtaaaaagaa ttataatgga attagacatg caaagcattc aagacattat tgctaaaaaa 60
ttaagtgggt ggcagaagag aaaactaaca ctagggattg ccatcttagg agatcctcag 120

<210> 102
<211> 139
<212> DNA
<213> Homo sapiens

<400> 102
gttttgctgc tagatgaacc aactgctgga ttggatccct tttcaagaca ccgagtgtgg 60
agcctcctga aggagcataa agtagaccga cttatcctct tcagtaccca attcatggat 120
gaggctgaca tcttggtg 139

<210> 103
<211> 91
<212> DNA
<213> Homo sapiens

<400> 103
ataggaaagt atttctgtct aatgggaagt tgaaatgtgc aggatcatct ttgtttctga 60
agcgaaagtg gggatttga tatcatttaa g 91

<210> 104
<211> 140
<212> DNA
<213> Homo sapiens

<400> 104
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tcctgatgcc aagttaacaa cagaaagtga agaaaaactt gtatatagtt tgccitttga 120
aaaaacgaac aaatttccag 140

<210> 105
<211> 120
<212> DNA
<213> Homo sapiens

<400> 105
atctttacag tgaccttgat aagtgttctg accagggcat aaggaattat gctgtttcag 60
tgacatctct gaatgaagta ttcttgaacc tagaaggaaa atcagcaatt gatgaaccag 120

<210> 106
<211> 199
<212> DNA
<213> Homo sapiens

<400> 106
atthttgacat tgggaaacaa gagaaaatac atgtgacaag aaatactgga gatgagtctg 60
aaatggaaca ggttctttgt tctcttcctg aaacaagaaa ggctgtcagt agtgcagctc 120
tctggagacg acaaattctat gcagtggcaa cacttcgctt cttaaagtta aggcgtgaaa 180
ggagagctct tttgtgttt 199

<210> 107
<211> 167
<212> DNA
<213> Homo sapiens

<400> 107
gttactagta cttggaattg cttttatccc catcattcta gagaagataa tgtataaagt 60
aactcgtgaa actcattgtt gggagttttc acccagtatg tatttccttt ctctggaaca 120
aatcccgaag acgcctctta ccagcctggt aatcgtaaat aatacag 167

<210> 108
<211> 134
<212> DNA
<213> Homo sapiens

<400> 108
gatcaaatat tgaagacctc gtgcattcac tgaagtgtca ggatatagtt ttggaaatag 60
atgactttag aaacagaaat ggctcagatg atccctccta caatggagcc atcatagtgt 120

ctggtgacca gaag

134

<210> 109
<211> 138
<212> DNA
<213> Homo sapiens

<400> 109
gattacagat tttctgttgc gtgtaatacc aagaaattga attgttttcc tgttcttatg 60
ggaattgtta gcaatgccct tatgggaatt ttttaacttca cggagcttat tcaaacggag 120
agcacttcat tttctcgt 138

<210> 110
<211> 108
<212> DNA
<213> Homo sapiens

<400> 110
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aactgcggtt ctccttttat cggcatgagc agcatcagcg attataaa 108

<210> 111
<211> 171
<212> DNA
<213> Homo sapiens

<400> 111
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caggctcttg tggacattcc attatacttc ttgattctct tttcaatata ttttaatttac 120
tacttcatat ttctgggatt ccagctttca tgggaactca tgtttgtttt g 171

<210> 112
<211> 114
<212> DNA
<213> Homo sapiens

<400> 112
gtggtatgca taattggttg tgcagtttct cttatattcc tcacatatgt gctttcattc 60
atctttcgca agtggagaaa aaataatggc ttttggctct ttggcttttt tatt 114

<210> 113
<211> 120
<212> DNA
<213> Homo sapiens

<400> 113
atcttaatat gtgtatccac aattatggta tcaactcaat atgaaaaact caacttaatt 60
ttgtgcatga ttttcatacc ttccttcact ttgctggggg atgtcatgtt attgatccag 120

<210> 114
<211> 81
<212> DNA
<213> Homo sapiens

<400> 114
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attcttttaa caaccttaat a 81

<210> 115
<211> 92
<212> DNA
<213> Homo sapiens

<400> 115
ccataccttc agagtgttat tttccttttt gtcataaggt gtctggaaat gaagtatgga 60
aatgaaataa tgaataaaga cccagttttc ag 92

<210> 116
<211> 121
<212> DNA
<213> Homo sapiens

<400> 116
aatctctcca cggagtagag aaactcatcc caatccggaa gagcccgaag aagaagatga 60
agatgttcaa gctgaaagag tccaagcagc aaatgcactc actgctccaa acttggagga 120
g 121

<210> 117
<211> 118
<212> DNA
<213> Homo sapiens

<400> 117
gaaccagtca taactgcaag ctgtttacac aaggaatatt atgagacaaa gaaaagttgc 60
ttttcaacaa gaaagaagaa aatagccatc agaaatgttt ctttttgtgt taaaaaag 118

<210> 118
<211> 92
<212> DNA
<213> Homo sapiens

<400> 118
gtgaagtttt gggattacta ggacacaatg gagctggtaa aagtacttcc attaaaatga 60
taactgggtg caciaagcca actgcaggag tg 92

<210> 119
<211> 179
<212> DNA
<213> Homo sapiens

<400> 119
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ttgggggtact gccctcagga gaactcactg tggcccaagc ttacaatgaa agagcacttg 120
gagttgtatg cagctgtgaa aggactgggc aaagaagatg ctgctctcag tatttcacg 179

<210> 120
<211> 76
<212> DNA
<213> Homo sapiens

<400> 120
attggtggaa gctcttaagc tccaggaaca acttaaggct cctgtgaaaa ctctatcaga 60
gggaataaag agaaag 76

<210> 121

<211> 95
 <212> DNA
 <213> Homo sapiens

<400> 121
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 accgggatgg accccgaggg gcagcagcaa atgtg 95

<210> 122
 <211> 120
 <212> DNA
 <213> Homo sapiens

<400> 122
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 catgtcagag gctgaggctg tgtgtgaccg tatggccatg atggtgtcag gaacgctaag 120

<210> 123
 <211> 141
 <212> DNA
 <213> Homo sapiens

<400> 123
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 aaaaatgaaa gaacctaccc aggtggaagc tctccacaca gagattttga agcttttccc 120
 acaggctgct tggcaggaaa g 141

<210> 124
 <211> 80
 <212> DNA
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<400> 124
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 ctttttcaag ttagaggcga 80

<210> 125
 <211> 56
 <212> DNA
 <213> Homo sapiens

<400> 125
 tgaaacagac cttcaacctg gaggaataca gcctctctca ggctaccttg gagcag 56

<210> 126
 <211> 769
 <212> DNA
 <213> Homo sapiens

<400> 126
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 acagttgaat ggaaacttct cccacaggaa gacccttaaa atgaagaacc tcctaacatt 120
 caatttttag tcctactaca ttgttagttt ccataattct acaagaatgt ttcctttttac 180
 ttcagttaac aaaagaaaac atttaataaa cattcaataa tgattacagt tttcattttt 240
 aaaaatttag gatgaaggaa acaaggaaat atagggaaaa gtagtagaca aaattaacaa 300
 aatcagacat gttattcatc cccaacatgg gtctattttg tgcttaaaaa taatttaaaa 360
 atcatacaat attaggtttg ttttcggtta ttatcaataa agctaactact gagaacattt 420
 tacaataaaa aatatgagtt ttttagcctg aacttcaaat gtatcagcta tttttaaaca 480
 ttattttactc ggatttcta ttaatgtgac attgactata agaaggctctg ataaactgat 540

gaaatggcac	agcataacat	ttaattataa	tgacattctg	attataaaat	aaatgcatgt	600
gaatttttagt	acatatgtgaa	gttatatgga	agaagatagc	cataatctgt	aagaaagtac	660
cgcagttaat	attttcttta	gccaaacttat	attcaatgta	ttttttatgg	atcctttttc	720
aaaggtagta	tcagtaggca	tagtcatttt	ctgtatcttt	tcacctcac		769

<210> 127
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<400> 127	
cagtactat	gtatccgtg
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gatggtttct	cctcacaac
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caccagacaa	tgaggatga
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gctatattct	tcaatggca
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cctagaagta	gaccgcctt
	19

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gttgtgagga	gaaaccatc
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gtacatgaaa actcaccata tccatccc	28
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gaaaatcagt ggcactcaat tc	22
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 <210> 180
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 actgtgaaag aaaacctcag gc 22

 <210> 181
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 aagaagaaat ggggcatagg 20

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 <210> 187
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 <210> 189
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 <210> 190
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